DATE: 05/15/2001

TIME: 17:26:37

PATENT APPLICATION: US/09/740,288A Input Set : A:\Pto.amc Output Set: N:\CRF3\05152001\1740288A.raw 3 <110> APPLICANT: Allen, Stephen Kinney, Anthony Miao, Guo-Hua 5 Orozco, Emil 6 8 <120> TITLE OF INVENTION: PLANT BIOTIN SYNTHASE 10 <130> FILE REFERENCE: BB1429 US NA 12 <140> CURRENT APPLICATION NUMBER: US 09/740288A 13 <141> CURRENT FILING DATE: 2000-12-19 15 <150> PRIOR APPLICATION NUMBER: US 60/172929 16 <151> PRIOR FILING DATE: 1999-12-21 18 <160> NUMBER OF SEQ ID NOS: 36 20 <170> SOFTWARE: Microsoft Office 97 23 <210> SEQ ID NO: 1 24 <211> LENGTH: 512 25 <212> TYPE: DNA 26 <213> ORGANISM: Hordeum vulgare 28 <220> FEATURE: 29 <221> NAME/KEY: Unsure 30 <222> LOCATION: (94)..(94) 31 <223> OTHER INFORMATION: n = A, C, G, or T34 <220> FEATURE: 35 <221> NAME/KEY: Unsure 36 <222> LOCATION: (460)..(460) 37 <223> OTHER INFORMATION: n = A, C, G, or T 40 <220> FEATURE: 41 <221> NAME/KEY: Unsure 42 <222> LOCATION: (462)..(462) 43 <223> OTHER INFORMATION: n = A, C, G, or T46 <400> SEQUENCE: 1 47 caactccctc ggcagtatcg cctagtgcag cagcggctcc gttccggcca gctttgctcg 60 W- $_{\mathcal{K}}>$  48 cegageegge catgatgetg etgetegege geanettege tecegegtee ggteeceett 120 49 egecteegee gttagegeeg egecettete ateggtateg geggeegegg eggaggegga 180 50 egggeggtge gggaegggee eaggaaegae tggaeeegee eegagateea ggeeatetae 240 51 gacteceege teetegacet cetetteeae ggggeteaag teeataggaa tgteeataaa 300 <del>52</del> tttagagaag tgcaacaatg cacacttctt tcaataaaga ctggtgggtg cagcgaagat 360 53 tgttcatact gcccacagtc ttcaagatac agtaccggat, tgaaggctga aaaattaatg 420 480 54 aagaaagatg ccgtcctaga agcagctaaa aaggcaaagn angctgggag cacccgattt 512 55 tgattggagc gatggagaga gacaattggc ag 58 <210> SEQ ID NO: 2 59 <211> LENGTH: 137 60 <212> TYPE: PRT 61 <213> ORGANISM: Hordeum vulgare 63 <220> FEATURE: 64 <221> NAME/KEY: UNSURE 65 <222> LOCATION: (131)..(131) 66 <223> OTHER INFORMATION: Xaa = any amino acid 69 <400> SEQUENCE: 2

RAW SEQUENCE LISTING

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/740,288A TIME: 05/15/2001
Input Set: A:\Pto.amc
Output Set: N:\CRF3\05152001\1740288A.raw

70 Met Met Leu Leu Leu Ala Arg Ser Leu Arg Ser Arg Val Arg Ser Pro 10 71 1 72 Phe Ala Ser Ala Val Ser Ala Ala Pro Phe Ser Ser Val Ser Ala Ala 73 20 74 Ala Ala Glu Ala Glu Arg Ala Val Arg Asp Gly Pro Arg Asn Asp Trp 75 35 40 76 Thr Arg Pro Glu Ile Gln Ala Ile Tyr Asp Ser Pro Leu Leu Asp Leu 78 Leu Phe His Gly Ala Gln Val His Arg Asn Val His Lys Phe Arg Glu 70 80 Val Gln Gln Cys Thr Leu Leu Ser Ile Lys Thr Gly Gly Cys Ser Glu 90 82 Asp Cys Ser Tyr Cys Pro Gln Ser Ser Arg Tyr Ser Thr Gly Leu Lys 105 100 110 84 Ala Glu Lys Leu Met Lys Lys Asp Ala Val Leu Glu Ala Ala Lys Lys 115 120 86 Ala Lys Xaa Ala Gly Ser Thr Arg Phe 130 90 <210> SEQ ID NO: 3 91 <211> LENGTH: 496 92 <212> TYPE: DNA 93 <213> ORGANISM: Zea mays 95 <220> FEATURE: 96 <221> NAME/KEY: Unsure 97 <222> LOCATION: (33)..(33) 98 <223> OTHER INFORMATION: n = A, C, G, or T 101 <220> FEATURE: 102 <221> NAME/KEY: Unsure 103 <222> LOCATION: (318)..(318) 104 <223> OTHER INFORMATION: n = A, C, G, or T 107 <220> FEATURE: 108 <221> NAME/KEY: Unsure 109 <222> LOCATION: (321)..(321) 110 <223> OTHER INFORMATION: n = A, C, G, or T 113 <220> FEATURE: 114 <221> NAME/KEY: Unsure 115 <222> LOCATION: (365)..(365) 116 <223> OTHER INFORMATION: n = A, C, G, or T 119 <220> FEATURE: 120 <221> NAME/KEY: Unsure 121 <222> LOCATION: (446)..(446) 122 <223> OTHER INFORMATION: n = A, C, G, or T 125 <400> SEQUENCE: 3 126 tccaatcggg tgggcagttt ttaaggaaac canggaccgc aagcaagcaa gccgccccag 127 ccgacgaggc gaggagcgtg caattccgta gctgcaacga actccctcga ccgtatcgcc

128 cgctgctcct ctatcccttt cctgctgctg ctactacctt aagctatcac tatcatggcc

129 ttgatgctgc tagcgcgcaa cctgcgctcc cgcctccgcc caccgctcgc cgccgccgcg

130 gqqttctcqt cqqccqcqqc qgagqcggag agggcgatac gggacgggcc gcggaacgac

120

180

240

300

360

DATE: 05/15/2001

TIME: 17:26:37

Input Set : A:\Pto.amc Output Set: N:\CRF3\05152001\1740288A.raw 132 ggggntcagt catcaagata caacactgga ttgaagggcc aaaaattgat gaacaaatat 480 🗲 133 gctgtcttgg gagcagcaaa aaaggnaaaa gagtctggga agcaaccgtt tttgcatggg 134 aactgcattg gagaaa 496 137 <210> SEQ ID NO: 4 138 <211> LENGTH: 102 139 <212> TYPE: PRT 140 <213> ORGANISM: Zea mays 142 <220> FEATURE: 143 <221> NAME/KEY: UNSURE 144 <222> LOCATION: (48)..(49) 145 <223> OTHER INFORMATION: Xaa = any amino acid 148 <220> FEATURE: 149 <221> NAME/KEY: UNSURE 150 <222> LOCATION: (64)..(64) 151 <223> OTHER INFORMATION: Xaa = any amino acid 154 <220> FEATURE: 155 <221> NAME/KEY: UNSURE 156 <222> LOCATION: (91)..(91) 157 <223> OTHER INFORMATION: Xaa = any amino acid 160 <400> SEQUENCE: 4 161 Met Ala Leu Met Leu Leu Ala Arg Asn Leu Arg Ser Arg Leu Arg Pro 163 Pro Leu Ala Ala Ala Gly Phe Ser Ser Ala Ala Glu Ala Glu 164 20 25 165 Arg Ala Ile Arg Asp Gly Pro Arg Asn Asp Trp Ser Arg Pro Glu Xaa 166 35 40 > 167 Xaa Ala Val Tyr Asp Ser Pro Leu Leu Asp Leu Leu Phe His Gly Xaa 168 55 169 Gln Ser Ser Arg Tyr Asn Thr Gly Leu Lys Gly Gln Lys Leu Met Asn 70 75~ 171 Lys Tyr Ala Val Leu Gly Ala Ala Lys Lys Xaa Lys Glu Ser Gly Lys 173 Gln Pro Phe Leu His Gly 174 100 177 <210> SEQ ID NO: 5 178 <211> LENGTH: 497 179 <212> TYPE: DNA 180 <213> ORGANISM: Zea mays 182 <220> FEATURE: 183 <221> NAME/KEY: Unsure 184 <222> LOCATION: (192)..(192) 185  $\langle 223 \rangle$  OTHER INFORMATION: n = A, C, G, or T 188 <220> FEATURE: 189 <221> NAME/KEY: Unsure 190 <222> LOCATION: (460)..(460) 191 <223> OTHER INFORMATION: n = A, C, G, or T 194 <220> FEATURE: 195 <221> NAME/KEY: Unsure 196 <222> LOCATION: (463)..(463)

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/740,288A

DATE: 05/15/2001

TIME: 17:26:37

Input Set : A:\Pto.amc Output Set: N:\CRF3\05152001\1740288A.raw 197 <223> OTHER INFORMATION: n = A, C, G, or T200 <220> FEATURE: 201 <221> NAME/KEY: Unsure 202 <222> LOCATION: (469)..(469) 203 <223> OTHER INFORMATION: n = A, C, G, or T206 <220> FEATURE: 207 <221> NAME/KEY: Unsure 208 <222> LOCATION: (490)..(490) 209 <223> OTHER INFORMATION: n = A, C, G, or T 212 <400> SEQUENCE: 5 213 ageegaegag gegaggageg tgeaatteeg tagetgeaae tgeaaegaae teeeteeete 214 octogacogt atogocogot gotoctotat coetttootg otgotgotac tacottaago 120 215 tatcatggcc ttgatgctgc tagcgcgcaa cctgcgctcc cgcctccgcc caccgctcgc 180 240  $-\Omega$  216 egeegeegeg gngttetegt eggeegegge ggaggeggag agggegatae gggaegggee 300 218 cetettteae ggggeteaag tecacagaaa tgtecataaa tteaagagaa gtgeageaat 219 gcacactict ticaatcaag actggtggga tgcagtgaag attgtictia cigtectcaa 420 220 gtcatcaaag aatacaacac tgggattgaa gggcccaaan aanttgatna acaaaagatg 480 221 ctgtcttggn aacaaca 497 224 <210> SEQ ID NO: 6 225 <211> LENGTH: 98 226 <212> TYPE: PRT 227 <213> ORGANISM: Zea mays 229 <220> FEATURE: 230 <221> NAME/KEY: UNSURE 231 <222> LOCATION: (23)..(23) 232 <223> OTHER INFORMATION: Xaa = any amino acid 235 <220> FEATURE: 236 <221> NAME/KEY: UNSURE 237 <222> LOCATION: (72)..(72) 238 <223> OTHER INFORMATION: Xaa = any amino acid 241 <220> FEATURE: 242 <221> NAME/KEY: UNSURE 243 <222> LOCATION: (89)..(89) 244 <223> OTHER INFORMATION: Xaa = any amino acid 247 <400> SEQUENCE: 6 248 Met Ala Leu Met Leu Leu Ala Arg Asn Leu Arg Ser Arg Leu Arg Pro 249 1 5 10 15 🗡 250 Pro Leu Ala Ala Ala Ala Xaa Phe Ser Ser Ala Ala Ala Glu Ala Glu 20 25 252 Arg Ala Ile Arg Asp Gly Pro Arg Asn Asp Trp Ser Arg Pro Glu Ile 40 254 Gln Ala Val Tyr Asp Ser Pro Leu Leu Asp Leu Leu Phe His Gly Ala 255 50 55 256 Gln Val His Arg Asn Val His Xaa Ser Arg Glu Val Gln Gln Cys Thr 257 65 70 75 258 Leu Leu Ser Ile Lys Thr Gly Gly Xaa Ser Glu Asp Cys Ser Tyr Cys

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/740,288A

260 Pro Gln

RAW SEQUENCE LISTING DATE: 05/15/2001 PATENT APPLICATION: US/09/740,288A TIME: 17:26:37

Input Set : A:\Pto.amc.

Output Set: N:\CRF3\05152001\I740288A.raw

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266 <212> TYPE: DNA
267 <213> ORGANISM: Zea mays
269 <400> SEQUENCE: 7
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271 egecegetge tectetatee ettteetget getgetaeta cettaageta teactateat
                                                                          180
272 ggccttgatg ctgctagcgc gcaacctgcg ctcccgcctc cgcccaccgc tcgccgccgc
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273 cgcggcgttc tcgtcggccg cggcggaggc ggagagggcg atacgggacg ggccgcggaa
                                                                          300
274 cgactggage cggcccgaga tecaggccgt ctacgactca ccgctcctcg acctcctctt
275 tcacggggct caggtccaca gaaatgtcca taaattcaga gaagtgcagc aatgcacact
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276 tettteaate aagactggtg gatgeagtga agattgttet tactgteete agteateaag
                                                                          420
                                                                          480
277 atacaacact ggattgaagg cccaaaaatt gatgaacaaa tatgctgtct tggaagcagc
                                                                          540
278 aaaaaaqqca aaaqaqtctq qqaqcacccq tttttqcatq qgaqctqcat ggaqaqaaac
279 cattqqcaqq aaatcaaact tcaaccagat tcttgaatat gtcaaggaaa taaggggtat
                                                                          600
280 gggcatggag gtctgttgca cactaggcat gatagagaaa caacaagctg aagaactcaa
                                                                          660
                                                                          720
281 gaaggetgga ettacageat ataateataa eetagataca teaagagagt attateeeaa
                                                                          780
282 cattattacc acaagatcat atgatgatag actgcagact cttgagcatg tccgtgaagc
283 tggaataagc atctgctcag gtggaatcat tggtcttggt gaagcagagg aggaccgggt
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284 agggttgttg cataccctag ctaccttgcc tacacaccca gagagcgttc ctattaatgc
                                                                          900
                                                                          960
285 attggttgct gtaaaaggca cacctcttga ggaccagaag cctgtagaga tctgggaaat
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286 gatecquatq atequated eteqqateac gatgecaaag gcaatggtga ggetttcage
                                                                         1080
287 aggccgagta cggttctcga tgccagaaca agcgctgtgc ttcctcgctg gggccaactc
288 catecttgee ggegagaaac tteteacaac egcaaacaac gaetttgatg eggaecaage
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293 <211> LENGTH: 344
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295 <213> ORGANISM: Zea mays
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299 1
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301
                                    25
302 Arg Ala Ile Arg Asp Gly Pro Arg Asn Asp Trp Ser Arg Pro Glu Ile
303
            35
                                40
304 Gln Ala Val Tyr Asp Ser Pro Leu Leu Asp Leu Leu Phe His Gly Ala
                            55
306 Gln Val His Arg Asn Val His Lys Phe Arg Glu Val Gln Gln Cys Thr
                                             75
                        70
307 65
308 Leu Leu Ser Ile Lys Thr Gly Gly Cys Ser Glu Asp Cys Ser Tyr Cys
                                        90
310 Pro Gln Ser Ser Arg Tyr Asn Thr Gly Leu Lys Ala Gln Lys Leu Met
                                    105
                100
311
312 Asn Lys Tyr Ala Val Leu Glu Ala Ala Lys Lys Ala Lys Glu Ser Gly
                                120
314 Ser Thr Arg Phe Cys Met Gly Ala Ala Trp Arg Glu Thr Ile Gly Arg
                            135
316 Lys Ser Asn Phe Asn Gln Ile Leu Glu Tyr Val Lys Glu Ile Arg Gly
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## Please Note:

Use f n and/or Xaa have been detected in the Sequence Listing. Pl as revi w the Sequence Listing to ensure that a corresp nding explanation is presented in the <220> to <223> fields of each sequence which presents at least ne n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/740,288A

DATE: 05/15/2001 TIME: 17:26:38

Input Set : A:\Pto.amc

Output Set: N:\CRF3\05152001\I740288A.raw

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L:54 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:86 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:133 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:165 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:167 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:171 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L\!:\!220~M\!:\!341~W\!: (46) "n" or "Xaa" used, for SEQ ID#:5
L:221 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:258 \ M:341 \ W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:365 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:588 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
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L:615 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:1517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
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